

Product datasheet for SR310332

OriGene Technologies, Inc.

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MKS1 Human siRNA Oligo Duplex (Locus ID 54903)

Product data:

Product Type: siRNA Oligo Duplexes

Purity: HPLC purified

Quality Control: Tested by ESI-MS

Sequences: Available with shipment

Stability: One year from date of shipment when stored at -20°C.

of transfections: Approximately 330 transfections/2nmol in 24-well plate under optimized conditions (final

conc. 10 nM).

Note: Single siRNA duplex (10nmol) can be ordered.

RefSeq: NM 001165927, NM 001321268, NM 001321269, NM 001330397, NM 017777

UniProt ID: Q9NXB0

Synonyms: BBS13; JBTS28; MES; MKS; POC12

Components: MKS1 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 54903)

Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol

Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml

Summary: The protein encoded by this gene localizes to the basal body and is required for formation of

the primary cilium in ciliated epithelial cells. Mutations in this gene result in Meckel syndrome type 1 and in Bardet-Biedl syndrome type 13. Multiple transcript variants encoding different

isoforms have been found for this gene. [provided by RefSeq, Sep 2009]

Performance OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will

Guaranteed: provide at least 70% or more knockdown of the target mRNA when used at 10 nM

concentration by quantitative RT-PCR when the TYE-563 fluorescent transfection control duplex (cat# SR30002) indicates that >90% of the cells have been transfected and the HPRT

positive control (cat# SR30003) provides 90% knockdown efficiency.

For non-conforming siRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the siRNA kit. To arrange for a free replacement with newly designed duplexes, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled siRNA control (quantitative RT-PCR data

required).

